SEQUENCE LISTING

<110> GARCIA-CALVO, MARIA MARGARITA	
<120> NPC1L1 (NPC3) AND METHODS OF IDENTIFYING LIGANDS THEREOF	
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Leu Ser Leu Glu Ser Ser Met Ser Ile Thr Lys Ala Leu Leu Thr Arg 100 105 110

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Gln Arg Ser Phe Ala Glu Lys Ala Tyr Glu Ser Cys Ser Gln Val Arg 165 170 175

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Glu Ala Pro Lys Leu Pro His Lys His Lys Leu Ser Pro His Thr Ile 325 330 335

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Leu Thr Val Leu Ala Leu Ser Phe Ile Val Val Ile Ala Leu Ala Ala 355 360 365

Gly Leu Thr Phe Ile Glu Leu Thr Thr Asp Pro Val Glu Leu Trp Ser 370 375 380

Ala Pro Lys Ser Gln Ala Arg Lys Glu Lys Ser Phe His Asp Glu His 385 390 395 400

Phe Gly Pro Phe Phe Arg Thr Asn Gln Ile Phe Val Thr Ala Arg Asn 405 410 415

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Val Met Leu Leu Phe Leu Thr Leu Phe Gly Ala Asn Leu Tyr Leu Met 850 855 860

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Glu Ala Gly Met Asn Ala Thr Cys Ser Ser Ala Gly Cys Lys Ser Phe 915 920 925

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gcc ac Ala Ti 12					agt Ser 1210							3654
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- Val Pro Ala Ala Thr Leu Ala Val Gly Thr Met Cys Gly Val Tyr 180 185 190
- Gly Ser Ala Leu Cys Asn Ala Gln Arg Trp Leu Asn Phe Gln Gly Asp 195 200 205
- Thr Gly Asn Gly Leu Ala Pro Leu Asp Ile Thr Phe His Leu Leu Glu 210 215 220
- Pro Gly Gln Ala Val Gly Ser Gly Ile Gln Pro Leu Asn Glu Gly Val 225 230 235 240

Ala Arg Cys Asn Glu Ser Gln Gly Asp Asp Val Ala Thr Cys Ser Cys 245 250 255

Gln Asp Cys Ala Ala Ser Cys Pro Ala Ile Ala Arg Pro Gln Ala Leu 260 265 270

Asp Ser Thr Phe Tyr Leu Gly Gln Met Pro Gly Ser Leu Val Leu Ile 275 280 285

Ile Ile Leu Cys Ser Val Phe Ala Val Val Thr Ile Leu Leu Val Gly 290 295 300

Phe Arg Val Ala Pro Ala Arg Asp Lys Ser Lys Met Val Asp Pro Lys 305 310 315 320

Lys Gly Thr Ser Leu Ser Asp Lys Leu Ser Phe Ser Thr His Thr Leu 325 330 335

Leu Gly Gln Phe Phe Gln Gly Trp Gly Thr Trp Val Ala Ser Trp Pro 340 345 350

Leu Thr Ile Leu Val Leu Ser Val Ile Pro Val Val Ala Leu Ala Ala 355 360 365

Gly Leu Val Phe Thr Glu Leu Thr Thr Asp Pro Val Glu Leu Trp Ser 370 375 380

Ala Pro Asn Ser Gln Ala Arg Ser Glu Lys Ala Phe His Asp Gln His 385 390 395 400

Phe Gly Pro Phe Phe Arg Thr Asn Gln Val Ile Leu Thr Ala Pro Asn 405 410 415

Arg Ser Ser Tyr Arg Tyr Asp Ser Leu Leu Gly Pro Lys Asn Phe 420 425 430

Ser Gly Ile Leu Asp Leu Asp Leu Leu Leu Glu Leu Glu Leu Gln 435 440 445

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Ile Ser Leu Gln Asp Ile Cys Tyr Ala Pro Leu Asn Pro Asp Asn Thr 465 470 475 480

Ser Leu Tyr Asp Cys Cys Ile Asn Ser Leu Leu Gln Tyr Phe Gln Asn 485 490 495

Asn Arg Thr Leu Leu Leu Leu Thr Ala Asn Gln Thr Leu Met Gly Gln 500 505 510

Thr Ser Gln Val Asp Trp Lys Asp His Phe Leu Tyr Cys Ala Asn Ala 515 520 525

Pro Leu Thr Phe Lys Asp Gly Thr Ala Leu Ala Leu Ser Cys Met Ala 530 540

Asp Tyr Gly Ala Pro Val Phe Pro Phe Leu Ala Ile Gly Gly Tyr Lys 545 550 555 560

Gly Lys Asp Tyr Ser Glu Ala Glu Ala Leu Ile Met Thr Phe Ser Leu 565 570 575

Asn Asn Tyr Pro Ala Gly Asp Pro Arg Leu Ala Gln Ala Lys Leu Trp 580 585 590

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Gly Met Phe Gln Val Thr Phe Thr Ala Glu Arg Ser Leu Glu Asp Glu 610 615 620

Ile Asn Arg Thr Thr Ala Glu Asp Leu Pro Ile Phe Ala Thr Ser Tyr 625 630 635 640

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Val Ala Val Val Leu Gly Ala Val Met Ala Ala Met Gly Phe Phe Ser 675 680 685

Tyr Leu Gly Ile Arg Ser Ser Leu Val Ile Leu Gln Val Val Pro Phe 690 695 700

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Ser Phe Thr Gln Lys Ile Gln Tyr Ala Thr Glu Phe Pro Glu Gln Ser

- Tyr Leu Ala Ile Pro Ala Ser Ser Trp Val Asp Asp Phe Ile Asp Trp 945 950 955 960
- Leu Thr Pro Ser Ser Cys Cys Arg Leu Tyr Ile Ser Gly Pro Asn Lys 965 970 975
- Asp Lys Phe Cys Pro Ser Thr Val Asn Ser Leu Asn Cys Leu Lys Asn 980 985 990
- Cys Met Ser Ile Thr Met Gly Ser Val Arg Pro Ser Val Glu Gln Phe 995 1000 1005
- His Lys Tyr Leu Pro Trp Phe Leu Asn Asp Arg Pro Asn Ile Lys 1010 1015 1020
- Cys Pro Lys Gly Gly Leu Ala Ala Tyr Ser Thr Ser Val Asn Leu 1025 1030 1035
- Thr Ser Asp Gly Gln Val Leu Ala Ser Arg Phe Met Ala Tyr His 1040 1045 1050
- Lys Pro Leu Lys Asn Ser Gln Asp Tyr Thr Glu Ala Leu Arg Ala 1055 1060 1065
- Ala Arg Glu Leu Ala Ala Asn Ile Thr Ala Asp Leu Arg Lys Val 1070 1075 1080
- Pro Gly Thr Asp Pro Ala Phe Glu Val Phe Pro Tyr Thr Ile Thr 1085 1090 1095
- Asn Val Phe Tyr Glu Gln Tyr Leu Thr Ile Leu Pro Glu Gly Leu 1100 1105 1110
- Phe Met Leu Ser Leu Cys Leu Val Pro Thr Phe Ala Val Ser Cys 1115 1120 1125
- Leu Leu Cly Leu Asp Leu Arg Ser Gly Leu Leu Asn Leu Leu 1130 1135 1140
- Ser Ile Val Met Ile Leu Val Asp Thr Val Gly Phe Met Ala Leu 1145 1150 1155

Trp Asp Ile Ser Tyr Asn Ala Val Ser Leu Ile Asn Leu Val Ser Ala Val Gly Met Ser Val Glu Phe Val Ser His Ile Thr Arg Ser Phe Ala Ile Ser Thr Lys Pro Thr Trp Leu Glu Arg Ala Lys Glu Ala Thr Ile Ser Met Gly Ser Ala Val Phe Ala Gly Val Ala Met Thr Asn Leu Pro Gly Ile Leu Val Leu Gly Leu Ala Lys Ala Gln Leu Ile Gln Ile Phe Phe Phe Arg Leu Asn Leu Leu Ile Thr Leu Leu Gly Leu Leu His Gly Leu Val Phe Leu Pro Val Ile Leu Ser Tyr Val Gly Pro Asp Val Asn Pro Ala Leu Ala Leu Glu Gln Lys Arg Ala Glu Glu Ala Val Ala Ala Val Met Val Ala Ser Cys Pro Asn His Pro Ser Arg Val Ser Thr Ala Asp Asn Ile Tyr Val Asn His Ser Phe Glu Gly Ser Ile Lys Gly Ala Gly Ala Ile Ser Asn Phe Leu Pro Asn Asn Gly Arg Gln Phe

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<213> Rattus sp.

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<213> Rattus sp.

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gtg tca Val Ser															;	336
tgc ccg Cys Pro															:	384
tgc agc Cys Ser 130	Pro															432
cgg gac Arg Asp 145																480
caa cgc Gln Arg															!	528
atc cct Ile Pro	gca Ala	gct Ala 180	gcc Ala	tcg Ser	ctg Leu	gct Ala	gtg Val 185	ggc Gly	agc Ser	atg Met	tgt Cys	gga Gly 190	gtg Val	tat Tyr	!	576
ggc tct Gly Ser	gcc Ala 195	ctc Leu	tgc Cys	aat Asn	gct Ala	cag Gln 200	cgc Arg	tgg Trp	ctc Leu	aac Asn	ttc Phe 205	caa Gln	gga Gly	gac Asp	ı	624
aca ggg Thr Gly 210	Asn	ggc Gly	ctg Leu	gct Ala	ccg Pro 215	ctg Leu	gac Asp	atc Ile	acc Thr	ttc Phe 220	cac His	ctc Leu	ttg Leu	gag Glu	ı	672

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	gc aat gag ys Asn Glu 245							768
	gt gca gca ys Ala Ala 260							816
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	tg gct tcc al Ala Ser							960
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ctt ggc co Leu Gly A	gg ttc ttc rg Phe Phe 340	gag agc	tgg gga Trp Gly 345	aca agg Thr Arg	gtg gcc Val Ala	tca tgg Ser Trp 350	cca Pro	1056
Leu Thr V	tc ttg gca al Leu Ala 55	Leu Ser	ttc ata Phe Ile 360	gtt gtg Val Val	ata gcc Ile Ala 365	Leu Ser	gta Val	1104
	cc ttt ata hr Phe Ile							1152
gcc cct ac Ala Pro L 385	aa agc caa ys Ser Gln	gcc cgg Ala Arg 390	aaa gaa Lys Glu	aag gct Lys Ala 395	ttc cat Phe His	gac gag Asp Glu	cat His 400	1200
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	gc tac aag er Tyr Lys 420							1296
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aac Asn	cac His	aca Thr	ctc Leu 500	ctg Leu	ctg Leu	ctc Leu	aca Thr	gcc Ala 505	aat Asn	cag Gln	act Thr	ctg Leu	aat Asn 510	ggc Gly	cag Gln	1536
	tcc Ser															1584
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	tac Tyr															1680
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tgc Cys 865	aac Asn	atc Ile	agc Ser	gtg Val	ggg Gly 870	ctg Leu	gac Asp	cag Gln	gat Asp	ctg Leu 875	gct Ala	ctg Leu	ccc Pro	aag Lys	gat Asp 880	2640
tcc Ser	tac Tyr	ctg Leu	ata Ile	gac Asp 885	tac Tyr	ttc Phe	ctc Leu	ttt Phe	ctg Leu 890	aac Asn	cgg Arg	tac Tyr	ttg Leu	gaa Glu 895	gtg Val	2688

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tat gtg gct att gct gca tcc tcc tgg gta gat gac ttc atc gac tgg Tyr Val Ala Ile Ala Ala Ser Ser Trp Val Asp Asp Phe Ile Asp Trp 945 950 955 960	2880
ctg acc cca tcc tcc tcc tgc tgc cgc att tat acc cgt ggc ccc cat Leu Thr Pro Ser Ser Ser Cys Cys Arg Ile Tyr Thr Arg Gly Pro His 965 970 975	2928
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4002

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Thr Ser Leu Ser Asn Ile Ser Cys Leu Ser Asn Thr Pro Ala Arg His 50 55 60

Val Thr Gly Asp His Leu Ala Leu Leu Gln Arg Val Cys Pro Arg Leu 65 70 75 80

Tyr Asn Gly Pro Asn Asp Thr Tyr Ala Cys Cys Ser Thr Lys Gln Leu 85 90 95

Val Ser Leu Asp Ser Ser Leu Ser Ile Thr Lys Ala Leu Leu Thr Arg 100 105 110

Cys Pro Ala Cys Ser Glu Asn Phe Val Ser Ile His Cys His Asn Thr 115 120 125

Cys Ser Pro Asp Gln Ser Leu Phe Ile Asn Val Thr Arg Val Val Gln 130 135 140

Arg Asp Pro Gly Gln Leu Pro Ala Val Val Ala Tyr Glu Ala Phe Tyr 145 150 155 160

Gln Arg Ser Phe Ala Glu Lys Ala Tyr Glu Ser Cys Ser Arg Val Arg 165 170 175

Ile Pro Ala Ala Ala Ser Leu Ala Val Gly Ser Met Cys Gly Val Tyr 180 185 190

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Gln Asp Cys	Ala Ala 260	Ser Cys	Pro	Val 265	Ile	Pro	Pro	Pro	Pro 270	Ala	Leu
Arg Pro Ser 275			280					285			
Ile Ile Phe 290		295					300				
Leu Arg Val 305		310				315					320
Glu Ala Pro	325				330					335	
Leu Gly Arg	340			345					350		
Leu Thr Val			360					365			
Gly Leu Thr 370		375					380				
Ala Pro Lys 385		390				395					400
Phe Gly Pro	Phe Phe 405	Arg Thr	Asn	GIn	11e 410	rne	vaı	Tnr	ATA	Lys 415	Asn

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Glu	Arg 450	Leu	Arg	His	Leu	Gln 455	Val	Trp	Ser	His	Glu 460	Ala	Gln	Arg	Asn
Ile 465	Ser	Leu	Gln	Asp	Ile 470	Cys	Tyr	Ala	Pro	Leu 475	Asn	Pro	His	Asn	Thr 480
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Gly	Thr	Asp	Tyr	Ser 565	Glu	Ala	Glu	Ala	Leu 570	Ile	Ile	Thr	Phe	Ser 575	Ile
Asn	Asn	Tyr	Pro 580	Ala	Asp	Asp	Pro	Arg 585	Met	Ala	His	Ala	Lys 590	Leu	Trp
Glu	Glu	Ala 595	Phe	Leu	Lys	Glu	Met 600	Gln	Ser	Phe	Gln	Arg 605	Ser	Thr	Ala
Asp	Lys 610	Phe	Gln	Ile	Ala	Phe 615	Ser	Ala	Glu	Arg	Ser 620	Leu	Glu	Asp	Glu
Ile 625	Asn	Arg	Thr	Thr	Ile 630	Gln	Asp	Leu	Pro	Val 635	Phe	Ala	Ile	Ser	Tyr 640
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Trp Ser Arg Val Ala Val Asp Ser Lys Ala Thr Leu Gly Leu Gly Gly 665 Val Ala Val Val Leu Gly Ala Val Val Ala Met Gly Phe Tyr Ser Tyr Leu Gly Val Pro Ser Ser Leu Val Ile Ile Gln Val Val Pro Phe Leu Val Leu Ala Val Gly Ala Asp Asn Ile Phe Ile Phe Val Leu Glu 710 Tyr Gln Arg Leu Pro Arg Met Pro Gly Glu Gln Arg Glu Ala His Ile 730 Gly Arg Thr Leu Gly Ser Val Ala Pro Ser Met Leu Cys Ser Leu Ser Glu Ala Ile Cys Phe Phe Leu Gly Ala Leu Thr Ser Met Pro Ala Val Arg Thr Phe Ala Leu Thr Ser Gly Leu Ala Ile Ile Phe Asp Phe 775 Leu Leu Gln Met Thr Ala Phe Val Ala Leu Leu Ser Leu Asp Ser Lys 790 795 785 Arg Gln Glu Ala Ser Arg Pro Asp Val Val Cys Cys Phe Ser Ser Arg 805 810 Asn Leu Pro Pro Lys Gln Lys Glu Gly Leu Leu Cys Phe Phe 825 830 820

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Arg Lys Ile Tyr Thr Pro Phe Leu Leu His Arg Phe Ile Arg Pro Val

Val Leu Leu Phe Leu Val Leu Phe Gly Ala Asn Leu Tyr Leu Met

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840

855

835

850

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- Ser Leu Thr Gln Lys Ile Gln Tyr Ala Ser Glu Phe Pro Asn Gln Ser 930 935 940
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- Lys Asp Glu Phe Cys Pro Ser Thr Asp Thr Ser Phe Asn Cys Leu Lys 980 985 990
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- Phe His Lys Tyr Leu Pro Trp Phe Leu Asn Asp Thr Pro Asn Ile 1010 1015 1020
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- His Lys Pro Leu Arg Asn Ser Gln Asp Phe Thr Glu Ala Leu Arg 1055 1060 1065
- Ala Ser Arg Leu Leu Ala Ala Asn Ile Thr Ala Glu Leu Arg Lys 1070 1075 1080
- Val Pro Gly Thr Asp Pro Asn Phe Glu Val Phe Pro Tyr Thr Ile 1085 1090 1095
- Ser Asn Val Phe Tyr Gln Gln Tyr Leu Thr Val Leu Pro Glu Gly 1100 1105 1110

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Val	Trp 1160	Gly	Ile	Ser	Tyr	Asn 1165	Ala	Val	Ser	Leu	Ile 1170	Asn	Leu	Val
Thr	Ala 1175	Val	Gly	Met	Ser	Val 1180	Glu	Phe	Val	Ser	His 1185	Ile	Thr	Arg
Ser	Phe 1190	Ala	Val	Ser	Thr	Lys 1195	Pro	Thr	Arg	Leu	Glu 1200	Arg	Ala	Lys
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Gln	Leu 1235	Ile	Gln	Ile	Phe	Phe 1240	Phe	Arg	Leu	Asn	Leu 1245	Leu	Ile	Thr
Leu	Leu 1250	Gly	Leu	Leu	His	Gly 1255	Leu	Val	Phe	Leu	Pro 1260	Val	Val	Leu
Ser	Tyr 1265	Leu	Gly	Pro		Val 1270		Gln	Ala		Val 1275		Glu	Glu
Lys	Leu 1280	Ala	Thr	Glu	Ala	Ala 1285	Met	Val	Ser	Glu	Pro 1290	Ser	Cys	Pro
Gln	Tyr 1295	Pro	Phe	Pro	Ala	Asp 1300	Ala	Asn	Thr	Ser	Asp 1305	Tyr	Val	Asn
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                                                                        107
Ala Glu Ala Gly Leu Arg Gly Trp Leu Leu Trp Ala Leu Leu Arg
ttg gcc cag agt gag cct tac aca acc atc cac cag cct ggc tac tgc
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Leu Ala Gln Ser Glu Pro Tyr Thr Thr Ile His Gln Pro Gly Tyr Cys
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gcc ttc tat gac gaa tgt ggg aag aac cca gag ctg tct gga agc ctc
Ala Phe Tyr Asp Glu Cys Gly Lys Asn Pro Glu Leu Ser Gly Ser Leu
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gtg ct Val Le	g tcc eu Ser	gtg Val	ggg Gly 710	gct Ala	gat Asp	aac Asn	atc Ile	ttc Phe 715	atc Ile	ttt Phe	gtt Val	ctc Leu	gag Glu 720	tac Tyr	2219

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ctg Leu	ccc Pro	ccg Pro 820	cct Pro	ggc Gly	cag Gln	gga Gly	gag Glu 825	ggg Gly	ctc Leu	ctg Leu	ctt Leu	ggc Gly 830	ttc Phe	ttc Phe	caa Gln	25	55
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Lys Ile Thr Gly Asp His Leu Ile Leu Leu Gln Lys Ile Cys Pro Arg 65 70 75 80

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Lys Gly Thr Ser Leu Ser Asp Lys Leu Ser Phe Ser Thr His Thr Leu 325 330 335

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Leu Thr Ile Leu Val Leu Ser Val Ile Pro Val Val Ala Leu Ala Ala 355 360 365

Gly Leu Val Phe Thr Glu Leu Thr Thr Asp Pro Val Glu Leu Trp Ser 370 380

Ala Pro Asn Ser Gln Ala Arg Ser Glu Lys Ala Phe His Asp Gln His 385 390 395 400

Phe Gly Pro Phe Phe Arg Thr Asn Gln Val Ile Leu Thr Ala Pro Asn 405 410 415

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Ser Gly Ile Leu Asp Leu Asp Leu Leu Leu Glu Leu Glu Leu Gln 435 440 445

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- Tyr Leu Gly Ile Arg Ser Ser Leu Val Ile Leu Gln Val Val Pro Phe 690 695 700
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